

FIG. 1A

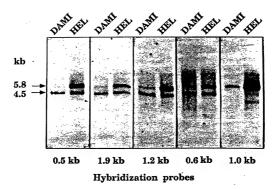
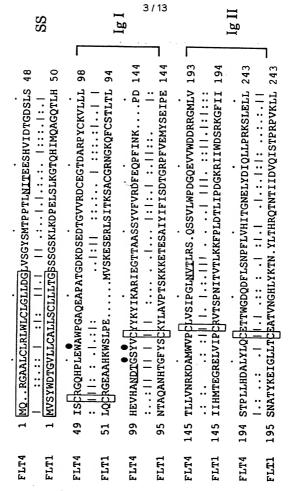


FIG. 1B



F16. 2A

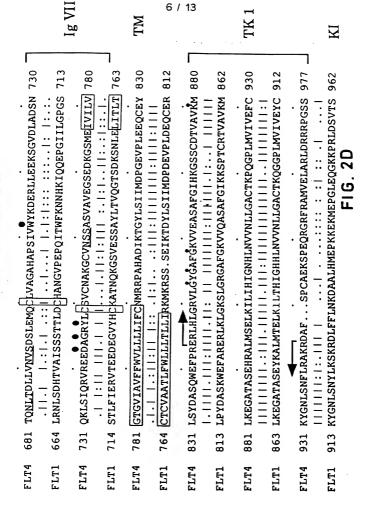
Ig	:	384 390 431 440	340 GPILEATAGDELVKLPVKLAAYPPPEFQWYKDGKALSGRHSPHAL 384 .::    . : .: : :  .  .     341 QQVLETVAGKRSYRLSMKVKAFPSPEVVWLKDGLPATEKSARYLTRGYSL 390 385 VLKEVTEASTGTYLALWNSAAGLRRNISLELVVNVPPQIHEKEASS 431 :: :    .  :   :  .    391 IIKDVTEEDAGNYTILLSIKQSNVFKNLTATLIVNVKPQIYEKAVSSFPD 440	340 341 385 391
	••••			
Ig	••••••	390	:    . : .:  :   .     ::  QQVLETVAGKRSYRLSMKVKAFPSPEVVWLKDGLPATEKSARXLTRGYSL	341
	•••••	384	GPILEATAGDELVKLPVKLAAYPPPEFQWYKDGKALSGRHSPHAL	340
	•••••	340	291 IFYSVLTIDKMQNKDKGLYTCRVRSGPSFKSVNTSVHIYDKAFITVKHRK 340	291
	٦			
70		339	ELSSILTIH <u>NVSQ</u> HDLGSYVEKANNGIQRFRESTEVIVHENPFISVEWLK 339	290
,		290	244 RGHTLVLNCHATTPLNTRVQMTWSYPDEKNRRASVRRIDQSNSHAN 290	244
		289	244 VGEKLVLNITTWABENSGVIFDWDYPGKQAERGKWVPERRSQQIHI 289	244

F16. 2B

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FLT4	432	432 PSIYSRHSROALTCTAYGVPLPLSIQWHWRPWTPCKMFAORSLRRROOOD 483	[	
FLT1	441	441 PALYPLGSRQILTCTAYGIPQP.TIKWFWHPCNHNHSEARCDFCSNNEES 489	101	
FLT4	482	482 LMPQCRDWRAVITQDAVNPIESLDTWTEFVEGK <u>NKT</u> VSKLVIQNANVSAM 531	531	
FLT1	490	FILDADSNMGNRIESITQRMAIIEGKNKMASTLVVADSRISGI	532	Ig V
FLT4	532	532 YKCVSNKVGQDERLIYFYVTTIPDGFTIESKPSEELLEGQPVLLSCDAD 581	.81	5 /
FLT1	533	533 YIÇIASNKVGTVGRNISFYITDVPNGFHVNLEKMPTEGEDLKLSCFVN 580	T 8	13
FLT4	582	582 SYKYEHLRWYRLN <u>LS</u> TLHDAHGNPLLLDCKNVHLFATPLAASLEEVAPGA 631	31	
FLT1	581	581 KFLYRDVTWILLRTVNNRTMHYSISKQKMAITK 613	13	Ig VI
FLT4	632	~	80	
FLT1	614	EHSITLNLTIMNVSLQDSGTYALRARNVYTGEEILQKKEITIRDQEAPYL 663	٦ وع	

F16. 2C

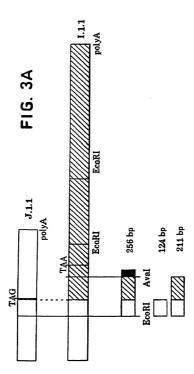


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LRDGTRMRAPELATPAIRRIMLNCWSGDPKARPAFSELVEILGDLLQGRG 1177_
                                                                                                                                                                                    1062
                                                                                                                                                                                                                                     LKWMAPESIFDKVYTTQSDVWSFGVLLWEIFSLGASPYPGVQINEEFCQR 1127
                                                                                                                                                                                                                                                                                                    FLT1 1063 LKWMAPESIFDKIYSTKSDVWSYGVLLWEIFSLGGSPYPGVQMDEDFCSR 1112
                                                                                                                                                                                                                                                                                                                                                                                                                      LREGMRMRAPEYSTPEIYQIMLDCWHRDPKERPRFAELVEKLGDLLQANV 1162
                                                                                                         FLT4 1028 LASRKCIHRDLAARNILLSESDVVKICDFGLARDIYKDPDYVRKGSARLP
                                                       963 SESFASSGFQEDKSLSDVEEEEDSDGFYKEPITMEDLISYSFQVARGMEF
978 DRVLFARFSKTEGGARRASPDQEAEDLWLSPLTMEDLVCYSFQVARGMEF
                                                                                                                                                                              FLT1 1013 LSSRKCIHRDLAARNILLSENNVVKICDFGLARDIYKNPDYVRKGDTRLP
                                                                                                                                                   F16. 2E
                                                                                                                                                                                                                                          FLT4 1078
                                                                                                                                                                                                                                                                                                                                                              1128
                                                                                                                                                                                                                                                                                                                                                                                                                      FLT1 1113
                                                                                                                                                                                                                                                                                                                                                                FLT4
  FLT4
                                                            FLT1
```

FLT4 1178 LQEEEEVCMAPRSSQSSEEGSFSQVSTMALHIAQADAEDSPPSLQRHSLA 1227 FLT1 1163 QQDGKDYIPINAILTGNSGFTYSTPAFSEDFFKESISAPKFNSGSSDDVR 1212 FLT1 1213 YVNAFKFMSLERIKTFEELLPNATSMFDDYQGDSSTLLASPMLKRFTWTD 1262 FLT4 1228 ARYYNWVSFPGCLARGAETRGSSRMKTFEEFPMTPTTYKGSVDNQTDSGM 1277 FLT4 1278 VLASEEFEQIESRHRQESGFR 1298

F16. 2F

FLT1 1263 SKPKASLKIEV 1273



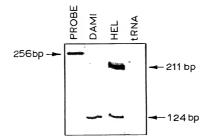


FIG. 3B

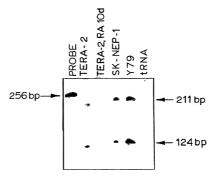


FIG. 3C

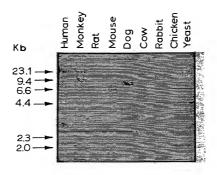


FIG. 4

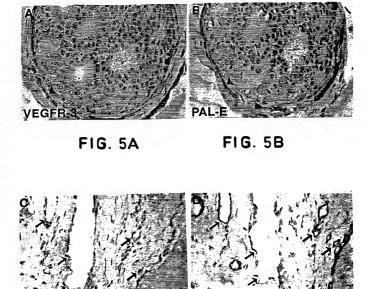
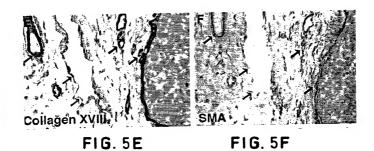


FIG. 5C

FIG. 5D

Laminin



VEGFR-3 and PAL-E/VEGFR-3
FIG. 5G
FIG. 5H